

SEQUENCE LISTING

```
<110> Yongjun Guo
      <120> SINGLE NUCLEOTIDE POLYMORPHISM IN THE
        FGF-3 GENE AND METHODS OF USE THEREOF
      <130> 3382-P03136US01
      <140> 10/798,652
      <141> 2004-03-11
      <150> 60/455,698
      <151> 2003-03-17
      <160> 7
      <170> FastSEQ for Windows Version 3.0
      <210> 1
      <211> 564
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (0)...(0)
      <223> n is cytosine or thymine
      <400> 1
gcagccctgc ctcagaaaac agaaggacgc agcacactca cggtgactca cccccatgtg
                                                                         60
                                                                        120
gctggaggng agggagcctc ctgaggcagg gccagggcag ccgtcaggtg ggtgacggca
ggggtcttgc catggtgggc acaggggctg catacagctt actcagtgac aatcgagtcc
                                                                        180
ctggtgccag cctctggaag tctggaagtg agcaatgttt cccattaagg aaagtgtgtg
                                                                        240
geoggecatg coeccaacg ttgcacacte actgcctttg cagggttggg gettecagte
                                                                        300
                                                                        360
acagggtccc atccacgtac cagcccaggt ggctgcagaa ggtccctcgc agtcatgaaa
ccaagggagg cttgggaaac cacatctgaa gggcatggct ttgatttagt gagagggtgg
                                                                        420
ggctgggctg ggcaaggcca ccaggtctga gtcagagcca gaggcaggaa gctggtcccc
                                                                        480
agcactgccc gccgcctctg cgatgcagtc ctcctggcca cctgagaaca gcctgtagag
                                                                        540
aggcagtggc gtctttcgga cttc
                                                                        564
      <210> 2
      <211> 20
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> primer
      <400> 2
gcagccctgc ctcagaaaac
                                                                         20
      <210> 3
      <211> 22
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> primer
      <400> 3
                                                                         22
tgcaccccac ttctagcatc ag
      <210> 4
      <211> 21
      <212> DNA
```

<213> Artificial Sequence	
<220>	
<223> primer	
<400> 4	
gcttcacccc agagatgagg g	21
<210> 5	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> primer	
<400> 5	
agctgtatgc agcccctgtg	20
2010.	
<210> 6	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> probe	
<400> 6	
ctccctcacc tccagccaca tg	22
<210> 7	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> probe	
/223/ brone	
<400> 7	
ctccctcgcc tccagccaca tg	22